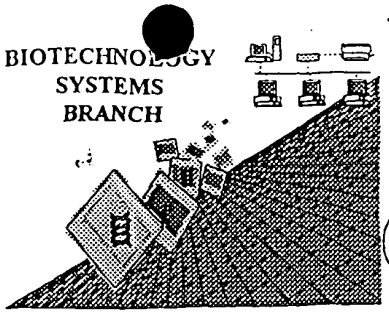
R. DeBerry

BIOTECHNOLOGY SYSTEMS BRANCH



Entered 7/19/01 RMD Paper 17

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/659,983B

Source: OIPE

Date Processed by STIC: 6/5/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.
Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

http://www.uspto.gov/web/offices/pac/checker

T. Debarry

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/659,983B

DATE: 06/05/2001
TIME: 14:22:06

Does Not Comply
Corrected Diskette Needed

Input Set : A:\PTO.txt
Output Set: C:\CRF3\06052001\I659983B.raw

pp 2,6

```
      3 <110> APPLICANT: Meloen, Robert Hans
      4       Oonk, Hendrica Berendina
      6 <120> TITLE OF INVENTION: An Improved Peptide, Immunogenic Composition and Vaccine or
Medical
      7       Preparation, a Method to Immunise Animals Against the Hormone LHRH, and Analogs
      8       of the LHRH Tandem Repeat Peptide and their Use as Vaccine
     10 <130> FILE REFERENCE: 2183-4518US
     12 <140> CURRENT APPLICATION NUMBER: 09/659,983B
C-> 13 <141> CURRENT FILING DATE: 2001-05-12
     15 <150> PRIOR APPLICATION NUMBER: US 09/274,048
     16 <151> PRIOR FILING DATE: 1999-03-22
     18 <150> PRIOR APPLICATION NUMBER: US 08/981,557
     19 <151> PRIOR FILING DATE: 1995-06-07
     21 <150> PRIOR APPLICATION NUMBER: PCT/NL96/00223
     22 <151> PRIOR FILING DATE: 1996-06-06
     24 <150> PRIOR APPLICATION NUMBER: US 08/447,298
     25 <151> PRIOR FILING DATE: 1995-06-07
     27 <150> PRIOR APPLICATION NUMBER: US 08/476,013
     28 <151> PRIOR FILING DATE: 1995-06-07
     30 <160> NUMBER OF SEQ ID NOS: 13
     32 <170> SOFTWARE: PatentIn version 3.0
     34 <210> SEQ ID NO: 1
     35 <211> LENGTH: 10
     36 <212> TYPE: PRT
     37 <213> ORGANISM: Sus scrofa
     39 <220> FEATURE:
     40 <221> NAME/KEY: PEPTIDE
     41 <222> LOCATION: (1)..(1)
     42 <223> OTHER INFORMATION: X=pyroglutamic acid
     45 <220> FEATURE:
     46 <221> NAME/KEY: PEPTIDE
     47 <222> LOCATION: (10)..(10)
     48 <223> OTHER INFORMATION: X=Gly-NH2
     51 <400> SEQUENCE: 1
W-> 53 Xaa His Trp Ser Tyr Gly Leu Arg Pro Xaa
     54 1               5                   10
     56 <210> SEQ ID NO: 2
     57 <211> LENGTH: 10
     58 <212> TYPE: PRT
     59 <213> ORGANISM: Homo sapiens
     61 <220> FEATURE:
     62 <221> NAME/KEY: PEPTIDE
     63 <222> LOCATION: (1)..(1)
     64 <223> OTHER INFORMATION: X=pyroglutamic acid
     67 <220> FEATURE:
     68 <221> NAME/KEY: PEPTIDE
     69 <222> LOCATION: (10)..(10)
     70 <223> OTHER INFORMATION: X=Gly-NH2
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ENTERED

RAW SEQUENCE LISTING DATE: 06/05/2001
PATENT APPLICATION: US/09/659,983B TIME: 14:22:06

Input Set : A:\PTO.txt
Output Set: C:\CRF3\06052001\I659983B.raw

73 <400> SEQUENCE: 2
W--> 75 **Xaa His Trp Ser His Gly Trp Tyr Pro Xaa**
76 1 5 10
78 <210> SEQ ID NO: 3
79 <211> LENGTH: 20
80 <212> TYPE: PRT
C--> 81 **<213> ORGANISM: artificial**
83 <220> FEATURE:
84 <223> OTHER INFORMATION: A peptide suitable for eliciting an immune response against forms
85 GnRH/ LHRH
87 <220> FEATURE:
88 <221> NAME/KEY: PEPTIDE
89 <222> LOCATION: (1)..(1)
90 <223> OTHER INFORMATION: X=pyroglutamic acid or Gln with attached tail of one or more addi
91 tional amino acid
94 <220> FEATURE:
95 <221> NAME/KEY: PEPTIDE
96 <222> LOCATION: (3)..(3)
97 <223> OTHER INFORMATION: X=Trp or N(indole)formyl-tryptophan
100 <220> FEATURE:
101 <221> NAME/KEY: PEPTIDE
102 <222> LOCATION: (11)..(11)
103 <223> OTHER INFORMATION: X=direct bond or a spacer group between Gly at position 10 and Gl
104 n at position 1
107 <220> FEATURE:
108 <221> NAME/KEY: PEPTIDE
109 <222> LOCATION: (13)..(13)
110 <223> OTHER INFORMATION: X=Trp or N(indole)formyl-tryptophan
113 <220> FEATURE:
114 <221> NAME/KEY: PEPTIDE
115 <222> LOCATION: (20)..(20)
116 <223> OTHER INFORMATION: X=Gly-NH2 or Gly with attached tail of one or more amino acids
119 <220> FEATURE:
120 <221> NAME/KEY: VARIANT
121 <222> LOCATION: (10)..(20)
122 <223> OTHER INFORMATION: variable repeat sequence <>10-20
125 <400> SEQUENCE: 3
W--> 127 **Xaa His Xaa Ser Tyr Gly Leu Arg Pro Gly Xaa His Xaa Ser Tyr Gly**
128 1 5 10 15
W--> 130 **Leu Arg Pro Xaa**
131 20
133 <210> SEQ ID NO: 4
134 <211> LENGTH: 21
135 <212> TYPE: PRT
C--> 136 **<213> ORGANISM: artificial**
138 <220> FEATURE:
139 <223> OTHER INFORMATION: A peptide suitable for eliciting an immune response against forms
140 GnRH/ LHRH

Xaa can only represent a single amino acid, nothing else

142 <220> FEATURE:

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/659,983B

DATE: 06/05/2001
TIME: 14:22:06

Input Set : A:\PTO.txt
Output Set: C:\CRF3\06052001\I659983B.raw

143 <221> NAME/KEY: PEPTIDE
144 <222> LOCATION: (1)..(1)
145 <223> OTHER INFORMATION: X=pyroglutamic acid
148 <220> FEATURE:
149 <221> NAME/KEY: PEPTIDE
150 <222> LOCATION: (6)..(6)
151 <223> OTHER INFORMATION: X=D-Lys
154 <220> FEATURE:
155 <221> NAME/KEY: PEPTIDE
156 <222> LOCATION: (11)..(11)
157 <223> OTHER INFORMATION: X=Gly or Gly preceded by a spacer
160 <220> FEATURE:
161 <221> NAME/KEY: PEPTIDE
162 <222> LOCATION: (16)..(16)
163 <223> OTHER INFORMATION: X=D-Lys
166 <220> FEATURE:
167 <221> NAME/KEY: PEPTIDE
168 <222> LOCATION: (21)..(21)
169 <223> OTHER INFORMATION: X=Cys-NH2
172 <400> SEQUENCE: 4
W--> 174 **Xaa His Thr Ser Tyr Xaa Leu Arg Pro Gly Xaa His Thr Ser Tyr Xaa**
175 1 5 10 15
W--> 177 **Leu Arg Pro Gly Xaa**
178 20
180 <210> SEQ ID NO: 5
181 <211> LENGTH: 21
182 <212> TYPE: PRT
183 <213> ORGANISM: artificial
185 <220> FEATURE:
186 <223> OTHER INFORMATION: A peptide suitable for eliciting an immune response against forms
187 GnRH/ LHRH
189 <220> FEATURE:
190 <221> NAME/KEY: PEPTIDE
191 <222> LOCATION: (1)..(1)
192 <223> OTHER INFORMATION: X=pyroglutamic acid
195 <220> FEATURE:
196 <221> NAME/KEY: PEPTIDE
197 <222> LOCATION: (4)..(4)
198 <223> OTHER INFORMATION: X=amino acid substitution
201 <220> FEATURE:
202 <221> NAME/KEY: PEPTIDE
203 <222> LOCATION: (6)..(6)
204 <223> OTHER INFORMATION: X=D-Lys
207 <220> FEATURE:
208 <221> NAME/KEY: PEPTIDE
209 <222> LOCATION: (11)..(11)
210 <223> OTHER INFORMATION: X=Gly or Gly preceded by a spacer
213 <220> FEATURE:
214 <221> NAME/KEY: PEPTIDE

RAW SEQUENCE LISTING DATE: 06/05/2001
PATENT APPLICATION: US/09/659,983B TIME: 14:22:06

Input Set : A:\PTO.txt
Output Set: C:\CRF3\06052001\I659983B.raw

215 <222> LOCATION: (14)..(14)
216 <223> OTHER INFORMATION: X=amino acid substitution
219 <220> FEATURE:
220 <221> NAME/KEY: PEPTIDE
221 <222> LOCATION: (16)..(16)
222 <223> OTHER INFORMATION: X=D-Lys
225 <220> FEATURE:
226 <221> NAME/KEY: PEPTIDE
227 <222> LOCATION: (21)..(21)
228 <223> OTHER INFORMATION: X=Cys-NH2
231 <400> SEQUENCE: 5
W--> 233 **Xaa His Thr Xaa Tyr Xaa Leu Ala Pro Gly Xaa His Thr Xaa Tyr Xaa**
234 1 5 10 15
W--> 236 **Leu Arg Pro Gly Xaa**
237 20
239 <210> SEQ ID NO: 6
240 <211> LENGTH: 21
241 <212> TYPE: PRT
OK> 242 **<213> ORGANISM: artificial**
244 <220> FEATURE:
245 <223> OTHER INFORMATION: A peptide suitable for eliciting an immune response against forms
246 GnRH/ LHRH
248 <220> FEATURE:
249 <221> NAME/KEY: PEPTIDE
250 <222> LOCATION: (1)..(1)
251 <223> OTHER INFORMATION: X=pyroglutamic acid
254 <220> FEATURE:
255 <221> NAME/KEY: PEPTIDE
256 <222> LOCATION: (6)..(6)
257 <223> OTHER INFORMATION: X=D-Lys
260 <220> FEATURE:
261 <221> NAME/KEY: PEPTIDE
262 <222> LOCATION: (8)..(8)
263 <223> OTHER INFORMATION: X=amino acid substitution
266 <220> FEATURE:
267 <221> NAME/KEY: PEPTIDE
268 <222> LOCATION: (11)..(11)
269 <223> OTHER INFORMATION: X=Gly or Gly preceded by a spacer
272 <220> FEATURE:
273 <221> NAME/KEY: PEPTIDE
274 <222> LOCATION: (16)..(16)
275 <223> OTHER INFORMATION: X=D-Lys
278 <220> FEATURE:
279 <221> NAME/KEY: PEPTIDE
280 <222> LOCATION: (18)..(18)
281 <223> OTHER INFORMATION: X=amino acid substitution
284 <220> FEATURE:
285 <221> NAME/KEY: PEPTIDE
286 <222> LOCATION: (21)..(21)

RAW SEQUENCE LISTING DATE: 06/05/2001
PATENT APPLICATION: US/09/659,983B TIME: 14:22:06

Input Set : A:\PTO.txt
Output Set: C:\CRF3\06052001\I659983B.raw

287 <223> OTHER INFORMATION: X=Cys-NH2
290 <400> SEQUENCE: 6
W--> 292 Xaa His Thr Ser Tyr Xaa Leu Xaa Pro Gly Xaa His Thr Ser Tyr Xaa
293 1 5 10 15
W--> 295 Leu Xaa Pro Gly Xaa
296 20
298 <210> SEQ ID NO: 7
299 <211> LENGTH: 21
300 <212> TYPE: PRT
W--> 301 <213> ORGANISM: artificial
303 <220> FEATURE:
304 <223> OTHER INFORMATION: A peptide suitable for eliciting an immune response against forms
305 GnRH/ LHRH
307 <220> FEATURE:
308 <221> NAME/KEY: PEPTIDE
309 <222> LOCATION: (1)..(1)
310 <223> OTHER INFORMATION: X=pyroglutamic acid
313 <220> FEATURE:
314 <221> NAME/KEY: PEPTIDE
315 <222> LOCATION: (6)..(6)
316 <223> OTHER INFORMATION: X=D-Lys
319 <220> FEATURE:
320 <221> NAME/KEY: PEPTIDE
321 <222> LOCATION: (10)..(10)
322 <223> OTHER INFORMATION: X=amino acid substutition
325 <220> FEATURE:
326 <221> NAME/KEY: PEPTIDE
327 <222> LOCATION: (11)..(11)
328 <223> OTHER INFORMATION: X=Gly or Gly preceded by a spacer
331 <220> FEATURE:
332 <221> NAME/KEY: PEPTIDE
333 <222> LOCATION: (16)..(16)
334 <223> OTHER INFORMATION: X=amino acid substitution
337 <220> FEATURE:
338 <221> NAME/KEY: PEPTIDE
339 <222> LOCATION: (20)..(20)
340 <223> OTHER INFORMATION: X=amino acid substitution
343 <220> FEATURE:
344 <221> NAME/KEY: PEPTIDE
345 <222> LOCATION: (21)..(21)
346 <223> OTHER INFORMATION: X=Cys-NH2
349 <400> SEQUENCE: 7
W--> 351 Xaa His Thr Ser Tyr Xaa Leu Arg Pro Xaa Xaa His Thr Ser Tyr Xaa
352 1 5 10 15
W--> 354 Leu Arg Pro Xaa Xaa
355 20
357 <210> SEQ ID NO: 8
358 <211> LENGTH: 42
359 <212> TYPE: PRT

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY DATE: 06/05/2001
PATENT APPLICATION: US/09/659,983B TIME: 14:22:07

Input Set : A:\PTO.txt
Output Set: C:\CRF3\06052001\I659983B.raw

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:53 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:75 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:81 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:127 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:130 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:136 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:174 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:177 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:183 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:233 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:236 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:242 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
L:292 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:295 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:301 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
L:351 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:354 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:360 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8
L:422 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:425 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:428 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:434 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
L:472 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:475 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:481 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10
L:519 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:522 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:528 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11
L:578 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:581 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:587 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12
L:637 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:640 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:646 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:13
L:690 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:693 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13